

Matches 17: Conservative 0; Mismatches 4; Indels 0; Gaps 0;

OY 5 tttgacatcttgcagca 25
|||||
Db 31 TGTGATCTTTATCAGCA 11

RESULT 6
A08043/c A08043 39 bp DNA PAT 05-AUG-1993
LOCUS Oligonucleotide P(450)11beta-6.
ACCESSION A08043
VERSION A08043.1 GI:413296
KEYWORDS
SOURCE Synthetic construct.
ORGANISM Synthetic construct
REFERENCE 1 (bases 1 to 39)
AUTHORS
JOURNAL Patent: WO 8910963-A 29-16-NOV-1989;
FEATURES
Location/Qualifiers
source 1..39
/organism="synthetic construct"
/db_xref="taxon:32630"

BASE COUNT 12 a 10 c 11 g 6 t
ORIGIN

Query Match 56.2%; Score 14.6; DB 6; Length 39;
Best Local Similarity 81.0%; Pred. No. 2.7e+04;
Matches 17: Conservative 0; Mismatches 4; Indels 0; Gaps 0;

OY 2 tattgtgacatcttgcaca 22
|||||
Db 27 TCTTGCCCATTTGTGCA 7

RESULT 7
AX008459 AX008459 20 bp DNA PAT 06-SEP-2000
LOCUS
DEFINITION Sequence 111 from Patent WO9966045.
ACCESSION AX008459
VERSION AX008459.1 GI:9996010
KEYWORDS
SOURCE Synthetic construct.
ORGANISM Synthetic construct
REFERENCE 1 (bases 1 to 20)
AUTHORS Gielkens, A.L., Koch, G., De Leeuw, O. and Peeters, B.P.
TITLE Newcastle disease virus infectious clones, vaccines and diagnostic assays
JOURNAL Patent: WO 9966045-A 111 23-DEC-1999;
DE (NL); PEETERS BERNARDUS PETRUS HUBER (NL); STICHTING DIENST
LANDBOUWKUNDE (NL)
FEATURES
Location/Qualifiers
source 1..20
/organism="synthetic construct"
/db_xref="taxon:32630"

primer_bind 1..20
/note="Primer P1898-, pos. 1898-1879,
primer"

BASE COUNT 4 a 4 c 6 g 6 t
ORIGIN

Query Match 55.4%; Score 14.4; DB 6; Length 20;
Best Local Similarity 93.8%; Pred. No. 3.3e+04;
Matches 15: Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 6 gtggacatcttgcac 21
|||||
Db 5 GTGGCCATCTTGTCC 20

RESULT 8
AR151074/c AR151074 39 bp DNA PAT 08-AUG-2001
LOCUS
DEFINITION Sequence 14 from patent US 6231850.
ACCESSION AR151074
VERSION AR151074.1 GI:15117124
KEYWORDS
SOURCE Unknown.
ORGANISM Unknown.
REFERENCE 1 (bases 1 to 39)
AUTHORS Okano, F., Satoh, M. and Yamada, K.
TITLE Canine interleukin 12
JOURNAL Patent: US 6231850-A 14 15-MAY-2001;
FEATURES
Location/Qualifiers
source 1..39
/organism="unknown"

BASE COUNT 10 a 14 c 8 g 7 t
ORIGIN

Query Match 54.6%; Score 14.2; DB 6; Length 39;
Best Local Similarity 84.2%; Pred. No. 4.1e+04;
Matches 16: Conservative 0; Mismatches 3; Indels 0; Gaps 0;

OY 7 tggacatcttgcagca 25
|||||
Db 33 TGGCATCTGTCTCTGCA 15

RESULT 9
E15028/c E15028 39 bp DNA PAT 28-JUL-1999
LOCUS
DEFINITION PCR primer for canine interleukin 12 40kDa subunit cDNA.
ACCESSION E15028
VERSION E15028.1 GI:5709711
KEYWORDS JP 1998036397-A/12.
SOURCE unidentified.
ORGANISM unidentified.
REFERENCE 1 (bases 1 to 39)
AUTHORS Okano, F.
TITLE CANINE INTERLEUKIN 12 AND ITS PRODUCTION
JOURNAL Patent: JP 1998036397-A 12 10-FEB-1998;
TORAY IND INC
OS None
COMMENT Artificial sequences.
PN JP 1998036397-A/12
PD 10-FEB-1998
PF 08-NOV-1996 JP 1996296789
PR 08-NOV-1995 JP 95P 289729, 23-MAY-1996 JP 96P 128104 PI
OKANO FUMIYOSHI
PC C07K14/54, C07H21/04, C12N5/10, C12N15/09, C12P21/02, (C12N5/10, PC
C12R1:91),
PC (C12P21/02, C12R1:91);
CC Strandedness: Single;
CC Topology: Linear;
FH key
FT source 1..39
Location/Qualifiers
source 1..39
/organism="unidentified"
/db_xref="taxon:32644"

BASE COUNT 10 a 14 c 8 g 7 t
ORIGIN

Query Match 54.6%; Score 14.2; DB 6; Length 39;
Best Local Similarity 84.2%; Pred. No. 4.1e+04;

Matches	16:	Conservative	0:	Mismatches	3:	Indels	0:	Gaps	0:
QY	7	tgggcatcttctgcacga	25						
Db	33	TTGGCATCTGTCTCTCCAG	15						
RESULT	10								
LOCUS	A67664	43 bp	DNA						
DEFINITION	Sequence 84 from Patent WO9744485.								
ACCESSION	A67664								
VERSION	A67664.1	GI:4756527							
KEYWORDS									
SOURCE	unidentified.								
ORGANISM	unclassified.								
REFERENCE	1 (bases 1 to 43)								
AUTHORS	Goodfellow,P.N.								
TITLE	METHODS FOR IDENTIFYING A MUTATION IN A GENE OF INTEREST								
JOURNAL	Patent: WO 97/44485-A 84 27-NOV-1997;								
FEATURES	HEXAGEN TECHNOLOGY LIMITED (GB)								
source	1..43								
BASE COUNT	12 a	14 c	7 g	10 t					
ORIGIN									
Query Match	54.6%;	Score 14.2;	DB 6;	Length 43;					
Best Local Similarity	84.2%;	Pred. No. 4.1e+04;							
Matches	16;	Conservative	0;	Mismatches	3;	Indels	0;	Gaps	0;
QY	5	ttggccatcttctgcacg	23						
Db	12	TATGACCATCTTCTCCAG	30						
RESULT	11								
LOCUS	AR089874	43 bp	DNA						
DEFINITION	Sequence 156 from patent US 5994075.								
ACCESSION	AR089874								
VERSION	AR089874.1	GI:10016629							
KEYWORDS									
SOURCE	Unknown.								
ORGANISM	Unknown.								
REFERENCE	1 (bases 1 to 43)								
AUTHORS	Goodfellow,P.N.								
TITLE	Methods for identifying a mutation in a gene of interest without a								
JOURNAL	phenotypic guide								
FEATURES	Patent: US 5994075-A 156 30-NOV-1999;								
source	1..43								
BASE COUNT	12 a	14 c	7 g	10 t					
ORIGIN									
Query Match	54.6%;	Score 14.2;	DB 6;	Length 43;					
Best Local Similarity	84.2%;	Pred. No. 4.1e+04;							
Matches	16;	Conservative	0;	Mismatches	3;	Indels	0;	Gaps	0;
QY	5	ttggccatcttctgcacg	23						
Db	12	TATGACCATCTTCTCCAG	30						
RESULT	12								
LOCUS	AX116937	51 bp	DNA						
ORIGIN									
Query Match	54.6%;	Score 14.2;	DB 6;	Length 43;					
Best Local Similarity	84.2%;	Pred. No. 4.1e+04;							
Matches	16;	Conservative	0;	Mismatches	3;	Indels	0;	Gaps	0;
QY	5	ttggccatcttctgcacg	23						
Db	12	TATGACCATCTTCTCCAG	30						
RESULT	12								
LOCUS	AX116937	51 bp	DNA						
ORIGIN									
Query Match	54.6%;	Score 14.2;	DB 6;	Length 43;					

DEFINITION	Sequence 2060 from Patent M00129262.
ACCESSION	AX116937
VERSION	AX116937.1
KEYWORDS	GI:14033879
SOURCE	human.
ORGANISM	Homo sapiens
REFERENCE	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
AUTHORS	Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.
TITLE	1 (bases 1 to 51)
JOURNAL	Picoult-Newburg, L. and Pohl, M.
	Genotyping reagents, kits and methods of use thereof
	Patent: WO 0129262-A 2060 26-APR-2001.
FEATURES	Orchid Biosciences, Inc. (US)
source	Location/Qualifiers
	1..51
	/organism="Homo sapiens"
	/db_xref="taxon:9606"
BASE COUNT	16 a 14 c 8 g 12 t 1 others
ORIGIN	
Query Match	54.6%; Score 14.2; DB 6; Length 51;
Best Local Similarity	84.2%; Pred. No. 4.2e+04;
Matches 16; Conservative 0; Mismatches 3; Indels 0; Gaps 0	
Oy 1 ttattgtgcatcttctgt 19	
11111111111111111111	
Db 35 TTGTGTGGCCATCTTAGT 17	
RESULT 13	
E64521/c	27 bp DNA PAT 07-FEB-2001
LOCUS	E64521
DEFINITION	Sugar-responsive enhancer in alpha-amylase gene.
ACCESSION	E64521
VERSION	E64521.1 GI:13017856
KEYWORDS	JP 1999332584-A/53.
SOURCE	unidentified.
ORGANISM	unclassified.
REFERENCE	1 (bases 1 to 27)
AUTHORS	Yoshimi, Y.
TITLE	Sugar-responsive enhancer in alpha-amylase gene
JOURNAL	Patent: JP 1999332584-A 53 07-DEC-1999;
COMMENT	NATIONAL SCIENCE COUNCIL
	OS Artificial Sequence
	PN JP 1999332584-A/53
	PD 07-DEC-1999
	PF 12-MAR-1999 JP 1999109867
	PR 12-MAR-1998 TW 87103633
	PI YOSHIMI YU
	PC C12N15/09,A01H1/00,C12N5/10,C12P21/02,C12N15/00,C12N5/00 CC
	Strandedness: Single;
	CC Topology: linear;
	FH key
	FT source
	1..27
	Location/Qualifiers
FEATURES	/organism='Artificial Sequence'.
source	1..27
	/organism="unidentified"
	/db_xref="taxon:32644"
BASE COUNT	12 a 7 c 5 g 3 t
ORIGIN	
Query Match	53.8%; Score 14; DB 6; Length 27;
Best Local Similarity	77.3%; Pred. No. 5.1e+04;
Matches 17; Conservative 0; Mismatches 5; Indels 0; Gaps 0;	
Oy 1 ttattgtgcatcttctgtcca 22	
11111111111111111111	
Db 26 TTATTGTGGTCCTCTATCGA 5	

RESULT 14

AX055762/c 40 bp DNA PAT 13-JAN-2001
 LOCUS AX055762
 DEFINITION Sequence 77 from Patent WO0073348.
 ACCESION AX055762
 VERSION AX055762.1 GI:12228874
 KEYWORDS

SOURCE synthetic construct.
 ORGANISM synthetic construct
 artificial sequence.

REFERENCE

1 (bases 1 to 40)
 Baker, K.P., Goddard, A., Gurney, A.L., Hebert, C., Henzel, W.,
 Kabakoff, R.C., Shelton, D.L., Smith, V., Watanabe, C.K. and Wood, W.I.
 Methods and compositions for inhibiting neoplastic cell growth
 Patent: WO 0073348-A 77 07-DEC-2000;
 Genentech, Inc. (US)

TITLE

Location/Qualifiers

FEATURES

1..40
 /organism="synthetic construct"
 /db_xref="taxon:32630"
 /note="Synthetic oligonucleotide probe."

BASE COUNT 10 a 9 c 12 g 9 t
 ORIGIN

Query Match

Best Local Similarity 77.3%; Score 14; DB 6; Length 40;
 Matches 17; Conservative 0; Mismatches 5; Indels 0; Gaps 0;

OY 5 ttatgtgcatcttgcacat 26
 ||| ||||| ||||| |||

Db 22 TGTCCCATCATCATCCACAT 1

RESULT 15

AR093976/c 41 bp DNA PAT 08-SEP-2000
 LOCUS AR093976
 DEFINITION Sequence 14 from patent US 6001595.
 ACCESION AR093976
 VERSION AR093976.1 GI:10020721
 KEYWORDS

SOURCE

Unknown.

ORGANISM

Unclassified.
 1 (bases 1 to 41)
 Ilmen, M., Onnela, M. and Penttila, M.
 Promoters and uses thereof
 Patent: US 6001595-A 14 14-DEC-1999;

REFERENCE

Location/Qualifiers

FEATURES

1..41
 /organism="unknown"

BASE COUNT

16 a 6 c 8 g 11 t

ORIGIN

Query Match 53.8%; Score 14; DB 6; Length 41;
 Best Local Similarity 77.3%; Pred. No. 5.1e+04;
 Matches 17; Conservative 0; Mismatches 5; Indels 0; Gaps 0;

OY 1 ttatgtgcatcttgcacat 22
 ||| ||||| ||||| |||

Db 25 TTAATGAGGCTATCTTATCGA 4

Search completed: March 9, 2002, 00:48:46
 Job time: 11127 sec

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